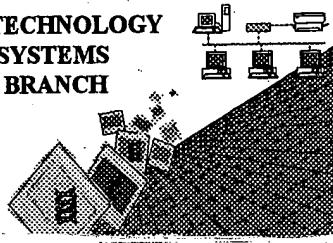


BIOTECHNOLOGY
SYSTEMS
BRANCH



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/057,475
 Source: OIPE
 Date Processed by STIC: 8/21/02

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FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

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Revised 01/29/2002



OIPE

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/057,475

DATE: 08/02/2002
TIME: 15:58:00

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Output Set: N:\CRF3\08022002\J057475.raw

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4     Algata, Paul A.
5     Mannion, Jane
6     Clapper, Jonathan David
7     Wang, Aijun
8     Ordonez, Nadia
9     Carter, Lauren
10    McNeill, Patricia Dianne
11    Corixa Corporation
13 <120> TITLE OF INVENTION: Compositions and Methods for the Detection, Diagnosis and
Therapy of
14    Hematological Malignancies
16 <130> FILE REFERENCE: 014058-014402US
18 <140> CURRENT APPLICATION NUMBER: US 10/057,475
C--> 19 <141> CURRENT FILING DATE: 2002-07-23
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22 <151> PRIOR FILING DATE: 2000-03-01
24 <150> PRIOR APPLICATION NUMBER: US 60/190,479
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RAW SEQUENCE LISTING
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/057,475

DATE: 08/02/2002
TIME: 15:58:42

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Output Set: N:\CRF3\08022002\J057475.raw

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PATENT APPLICATION: US/10/057,475

DATE: 08/02/2002
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RAW SEQUENCE LISTING
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W--> 214832	cnagnaccaa tctgggtctc agggatnata cccggcttta g	401
219976 <210>	SEQ ID NO: 10972	
219977 <211>	LENGTH: 471	
219978 <212>	TYPE: DNA	
219979 <213>	ORGANISM: Homo sapiens	some error
219981 <221>	NAME/KEY: unsure	
219982 <222>	LOCATION: (1)...(471)	
219983 <223>	OTHER INFORMATION: n = g, a, c or t	
E--> 219985 <400>	SEQUENCE: 10972	
219986	aaatttagga cacggtacat tttccactga aaatttacaa tcataccaaca taatgcacac	60
219987	caccctcaa aggtaacatt agctcatata caaaaatcatg gaaaacctca caacatccctt	120
219988	ggaaggtaga tattgttata tccttacaaa aatttaatac accattaaac attccttattt	180
W--> 219989	caaatagctc aatcgatatac aacacataag acagccccaca attgtgatac agnattaaga	240
219990	agaaactcag gccagcccg gtgcctcagc cctgttaaccc caacactttg ggaggctgag	300
219991	gcgggcagat cacctgagcc caggagtca ggaccagct ggacaacatg gtgagacccc	360
219992	gtctctacaa aaagtacaaa aattagctgg gtgtggtgac ccacgcctgt agtcccagct	420
W--> 219993	agttgggagg ctaagacggg aggatggctn gttgcttgag cccaaaggagg t	471
219995 <210>	SEQ ID NO: 10973	
219996 <211>	LENGTH: 379	
219997 <212>	TYPE: DNA	
219998 <213>	ORGANISM: Homo sapiens	some error
220000 <221>	NAME/KEY: unsure	
220001 <222>	LOCATION: (1)...(379)	
220002 <223>	OTHER INFORMATION: n = g, a, c or t	
E--> 220004 <400>	SEQUENCE: 10973	
W--> 220005	aaaataagct atggttttc cagtagccaa aatgatcctg caccanagct catanactga	60
W--> 220006	gaacctganc atgcaaanc acagtctggg tgaaggatg tctgtttgt aaatgacctg	120
220007	ctaattcttt gcaacccaca gtaatttggt ttctgtgaac ccacagaagc agggccacca	180
W--> 220008	aaaagggctc tgtctgctag cctggatcat acatgantca ctggcggtgg gatcagtcat	240
220009	ttttaggct gccccattt cctaacatgt taaaatgtgt gttctcagtc tttcaagag	300
W--> 220010	aggaagaagc aaagcggcac ttacagatgt tgtgatanga cacagatctg tggcgaggga	360
220011	ttgggaagg tgggtggca	379

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L:19 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:105 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:106 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:60
L:131 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
L:135 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:240
L:137 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:360
L:171 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:420
L:191 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:360
L:192 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:420
L:205 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:207 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:120
L:209 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:240
L:210 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:300
L:211 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:360
L:229 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:300
L:244 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:60
L:245 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:120
L:246 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:180
L:247 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:240
L:267 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:420
L:296 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:300
L:323 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:180
L:324 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:240
L:325 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:300
L:326 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:360
L:327 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:420
L:343 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:180
L:368 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
L:369 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:60
L:370 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:120
L:371 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:180
L:400 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:120
L:401 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:180
L:402 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:240
L:415 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:417 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:120
L:431 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:60
L:479 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:300
L:480 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:360
L:494 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
L:497 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:180
L:499 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:300
L:514 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:60
L:515 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:120
L:541 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0
L:543 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:120
L:544 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:180
L:547 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:360

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L:563 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:180
L:566 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:360
L:567 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:420
L:57901 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2533
L:57904 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2533
L:57908 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2533
L:57944 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2535
L:57947 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2535
L:57950 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2535
L:57953 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2535
L:57956 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2535
L:57959 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2535
L:57962 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2535
L:57965 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2535
L:57969 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2535
L:57988 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2536
L:57991 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2536
L:57994 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2536
L:57997 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2536
L:58000 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2536
L:58003 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2536
L:58006 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2536
L:58009 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2536
L:58013 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2536
L:58066 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2539
L:58104 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2541
L:58108 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2541
L:58128 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2542
L:58147 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2543
L:58150 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2543
L:58153 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2543
L:58156 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2543
L:58159 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2543
L:58163 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2543
L:58206 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2546
L:58285 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2551
L:58305 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2552
L:58347 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2555
L:58351 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2555
L:58405 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2558
L:58409 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2558
L:58445 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2560
L:58485 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2562
L:58489 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2562
L:58549 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2566
L:58640 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2572
L:58643 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2572
L:58646 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2572
L:58649 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2572

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L:58653 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2572
L:58673 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2573
L:58946 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2594
L:58990 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2597
L:211722 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:10475
L:214678 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:10581
L:214710 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:10583
L:214730 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:10584
L:214745 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:10585
L:214768 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:10586
L:214805 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:10588
L:214825 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:10589
L:217674 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (10819) SEQUENCE:
L:219985 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:10972
L:220004 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:10973
L:63 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (10969) Counted (10979)